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<110> ANDERSEN, Peter
      SKJOT, Rikke
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<213> Mycobacterium tuberculosis

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Glu Asp Tyr Gly Val Thr Ile Ala Asp Gly Pro Met Ala Gly Leu Leu 115 120 125

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Ala Ser Asp Asp Tyr Arg Ala Ser Ala Ser Asn Gly Ser Asp Asp Ala 85 90 95

Ser Ala His Ile Gln Arg Thr Val Ala Ser Cys Pro Asn Thr Arg Ile

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100 105 110

Val Leu Gly Gly Tyr Ser Gln Gly Ala Thr Val Ile Asp Leu Ser Thr
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Cys Ala Pro Asp Asp Pro Ile Cys Thr Gly Gly Gly Asn Ile Met Ala 180 185 190

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Gln Gly Gly Asp Pro Thr Gly Thr Gly Arg Gly Gly Pro Gly Tyr Lys 85 90 95

Phe Ala Asp Glu Phe His Pro Glu Leu Gln Phe Asp Lys Pro Tyr Leu 100 105 110

Leu Ala Met Ala Asn Ala Gly Pro Gly Thr Asn Gly Ser Gln Phe Phe 115 120 125

Ile Thr Val Gly Lys Thr Pro His Leu Asn Arg Arg His Thr Ile Phe 130 135 140

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- Val Ile Ala His Leu Arg Ala Ser Lys Pro Leu Val Arg Leu Arg Val
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- Pro Phe Thr Leu Ser Arg Asn Glu Ile Asp Asp Val Glu Arg Gly Ser 85 90 95
- Lys Asp Ser Asp Trp Glu Pro Val Lys Glu Ala Ala Lys Lys Leu Ala 100 105 110
- Phe Val Glu Asp Arg Thr Ile Phe Glu Gly Tyr Ser Ala Ala Ser Ile 115 120 125
- Glu Gly Ile Arg Ser Ala Ser Ser Asn Pro Ala Leu Thr Leu Pro Glu 130 135 140
- Asp Pro Arg Glu Ile Pro Asp Val Ile Ser Gln Ala Leu Ser Glu Leu 145 150 155 160
- Arg Leu Ala Gly Val Asp Gly Pro Tyr Ser Val Leu Leu Ser Ala Asp 165 170 175
- Val Tyr Thr Lys Val Ser Glu Thr Ser Asp His Gly Tyr Pro Ile Arg 180 185 190
- Glu His Leu Asn Arg Leu Val Asp Gly Asp Ile Ile Trp Ala Pro Ala 195 200 205
- Ile Asp Gly Ala Phe Val Leu Thr Thr Arg Gly Gly Asp Phe Asp Leu 210 215 220
- Gln Leu Gly Thr Asp Val Ala Ile Gly Tyr Ala Ser His Asp Thr Asp

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gcatgcagca attcggcggt gtggacacca acggaatgtg gggagcacca cagctgggtc 660
ggtggaagtg gcacgacccg tgggtgcatg ccagcctgct ggcgcaaaac aacacccggg 720
tgtgggtgtg gagcccgacc aacccgggag ccagcgatcc cgccgccatg atcggccaaa 780
ccgccgaggc gatgggtaac agccgcatgt tctacaacca gtatcgcagc gtcggcgggc 840
acaacggaca cttcgacttc ccagccagcg gtgacaacgg ctgggggctcg tgggcgcccc 900
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<210> 42
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<212> PRT

<213> Mycobacterium tuberculosis

<400> 42

Met Lys Gly Arg Ser Ala Leu Leu Arg Ala Leu Trp Ile Ala Ala Leu 1 5 10 15

Ser Phe Gly Leu Gly Gly Val Ala Val Ala Ala Glu Pro Thr Ala Lys 20 25 30

Ala Ala Pro Tyr Glu Asn Leu Met Val Pro Ser Pro Ser Met Gly Arg
35 40 45

Asp Ile Pro Val Ala Phe Leu Ala Gly Gly Pro His Ala Val Tyr Leu 50 55 60

Leu Asp Ala Phe Asn Ala Gly Pro Asp Val Ser Asn Trp Val Thr Ala 65 70 75 80

Gly Asn Ala Met Asn Thr Leu Ala Gly Lys Gly Ile Ser Val Val Ala 85 90 95

Pro Ala Gly Gly Ala Tyr Ser Met Tyr Thr Asn Trp Glu Gln Asp Gly
100 105 110

Ser Lys Gln Trp Asp Thr Phe Leu Ser Ala Glu Leu Pro Asp Trp Leu 115 120 125

Ala Ala Asn Arg Gly Leu Ala Pro Gly Gly His Ala Ala Val Gly Ala 130 135 140

Ala Gln Gly Gly Tyr Gly Ala Met Ala Leu Ala Ala Phe His Pro Asp 145 150 155 160

Arg Phe Gly Phe Ala Gly Ser Met Ser Gly Phe Leu Tyr Pro Ser Asn 165 170 175

Thr Thr Thr Asn Gly Ala Ile Ala Ala Gly Met Gln Gln Phe Gly Gly
180 185 190

Val Asp Thr Asn Gly Met Trp Gly Ala Pro Gln Leu Gly Arg Trp Lys 195 200 205

Trp His Asp Pro Trp Val His Ala Ser Leu Leu Ala Gln Asn Asn Thr 210 215 220

Arg Val Trp Val Trp Ser Pro Thr Asn Pro Gly Ala Ser Asp Pro Ala 225 230 235 240

Ala Met Ile	Gly Gln 245	Thr Ala	Glu Al	a Met 250	Gly	Asn	Ser	Arg	Met 255	Phe	
Tyr Asn Gln	Tyr Arg 260	Ser Val	Gly Gl		Asn	Gly	His	Phe 270	Asp	Phe	
Pro Ala Ser 275	Gly Asp	Asn Gly	Trp Gl 280	y Ser	Trp	Ala	Pro 285	Gln	Leu	Gly	
Ala Met Ser 290	Gly Asp	Ile Val 295	Gly Al	a Ile	Arg						
<210> 43 <211> 27 <212> DNA <213> Mycoba	acterium	tubercu	losis								
<400> 43 gcaacacccg g	ggatgtcgo	ca aatca	tg								27
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<400> 44 gtaacacccg g											27
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<210> 46 <211> 38	33340000	ou 30030		-55-5	9						
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<210> 47 <211> 450											

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<212> DNA
<213> Mycobacterium tuberculosis
<400> 47
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gcgccgagat cgtggccagc gttctcgaag tcgttgtcaa cgaaggcgat cagatcgaca 180
agggcgacgt cgtggtgctg ctggagtcga tgaagatgga gatccccgtc ctggccgaag 240
ctgccqqaac gqtcagcaag gtggcggtat cggtgggcga tgtcattcag gccggcgacc 300
ttatcgcggt gatcagctag tcqttgatag tcactcatgt ccacactcgg tgatctgctc 360
gccgaacaca cggtgctgcc gggcagcgcg gtggaccacc tgcatgcggt ggtcggggag 420
tggcagetee ttgccgaett gtcgtttgcc
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<211> 71
<212> PRT
<213> Mycobacterium tuberculosis
<400> 48
Met Ala Glu Asp Val Arg Ala Glu Ile Val Ala Ser Val Leu Glu Val
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Val Val Asn Glu Gly Asp Gln Ile Asp Lys Gly Asp Val Val Leu
                                 25
Leu Glu Ser Met Lys Met Glu Ile Pro Val Leu Ala Glu Ala Ala Gly
                             40
         35
Thr Val Ser Lys Val Ala Val Ser Val Gly Asp Val Ile Gln Ala Gly
     50
Asp Leu Ile Ala Val Ile Ser
 65
                     70
<210> 49
<211> 750
<212> DNA
<213> Mycobacterium tuberculosis
<400> 49
gggtacccat cgatgggttg cggttcggca ccgaggtgct aacgcacttg ctgacacact 60
gctagtcgaa aacgaggcta gtcgcaacgt cgatcacacg agaggactga ccatgacaac 120
ttcaccegac cegtatgeeg egetgeecaa getgeegtee ttcageetga egteaacete 180
gatcaccgat gggcagccgc tggctacacc ccaggtcagc gggatcatgg gtgcgggcgg 240
ggeggatgee agteegeage tgaggtggte gggattteee agegagaeee geagettege 300
ggtaaccgtc tacgaccctg atgcccccac cctgtccggg ttctggcact gggcggtggc 360
caacctgcct gccaacgtca ccgagttgcc cgagggtgtc ggcgatggcc gcgaactgcc 420
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gggcggggca ctgacattgg tcaacgacgc cggtatgcgc cggtatgtgg gtgcggcgcc 480 gcctcccggt catggggtgc atcgctacta cgtcgcggta cacgcggtga aggtcgaaaa 540 gctcgacctc cccgaggacg cgagtcctgc atatctggga ttcaacctgt tccagcacgc 600 gattgcacga gcggtcatct tcggcaccta cgagcagcgt tagcgcttta gctgggttgc 660 cgacgtcttg ccgagccgac cgcttcgtgc agcgagccga acccgccgtc atgcacctg 720 cgggcaatgc cttcatggat gtccttggcc

<210> 50

<211> 176

<212> PRT

<213> Mycobacterium tuberculosis

<400> 50

Met Thr Thr Ser Pro Asp Pro Tyr Ala Ala Leu Pro Lys Leu Pro Ser 1 5 10 15

Phe Ser Leu Thr Ser Thr Ser Ile Thr Asp Gly Gln Pro Leu Ala Thr 20 25 30

Pro Gln Val Ser Gly Ile Met Gly Ala Gly Gly Ala Asp Ala Ser Pro 35 40 45

Gln Leu Arg Trp Ser Gly Phe Pro Ser Glu Thr Arg Ser Phe Ala Val 50 55 60

Thr Val Tyr Asp Pro Asp Ala Pro Thr Leu Ser Gly Phe Trp His Trp 65 70 75 80

Ala Val Ala Asn Leu Pro Ala Asn Val Thr Glu Leu Pro Glu Gly Val 85 90 95

Gly Asp Gly Arg Glu Leu Pro Gly Gly Ala Leu Thr Leu Val Asn Asp 100 105 110

Ala Gly Met Arg Arg Tyr Val Gly Ala Ala Pro Pro Pro Gly His Gly
115 120 125

Val His Arg Tyr Tyr Val Ala Val His Ala Val Lys Val Glu Lys Leu 130 135 140

Asp Leu Pro Glu Asp Ala Ser Pro Ala Tyr Leu Gly Phe Asn Leu Phe 145 150 155 160

Gln His Ala Ile Ala Arg Ala Val Ile Phe Gly Thr Tyr Glu Gln Arg 165 170 175

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<210> 51
<211> 800
<212> DNA
<213> Mycobacterium tuberculosis
<400> 51
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cccgcggaac cggcgaacca cctggcctcg gtcgggtagg ccaagctttc gtcagttcat 240
tgcgccagca gaccaacaag agcatcggga catacggagt caactacccg gccaacggtg 300
atttcttggc cgccgctgac ggcgcgaacg acgccagcga ccacattcag cagatggcca 360
gegegtgeeg ggeeacgagg ttggtgeteg geggetaete ceagggtgeg geegtgateg 420
acategteac egeegeacea etgeeeggee tegggtteac geageegttg eegeeegeag 480
cggacgatca catcgccgcg atcgccctgt tcgggaatcc ctcgggccgc gctggcgggc 540
tgatgagege cetgacecet caattegggt ccaagaceat caacetetge aacaaeggeg 600
accegatttg ttcggacggc aaccggtggc gagcgcacct aggctacgtg cccgggatga 660
ccaaccagge ggegegttte gtegegagea ggatetaaeg egageegeee catagattee 720
ggctaagcaa cggctgcgcc gccgcccggc cacgagtgac cgccgccgac tggcacaccg 780
cttaccacgg ccttatgctg
<210> 52
<211> 226
<212> PRT
<213> Mycobacterium tuberculosis
<400> 52
Met Ile Pro Arg Pro Gln Pro His Ser Gly Arg Trp Arg Ala Gly Ala
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                                      10
                   5
  1
Ala Arg Arg Leu Thr Ser Leu Val Ala Ala Ala Phe Ala Ala Ala Thr
                                                      30
              20
                                  25
Leu Leu Leu Thr Pro Ala Leu Ala Pro Pro Ala Ser Ala Gly Cys Pro
                                                   45
                              40
          35
 Asp Ala Glu Val Val Phe Ala Arg Gly Thr Gly Glu Pro Pro Gly Leu
                                              60
      50
                          55
 Gly Arg Val Gly Gln Ala Phe Val Ser Ser Leu Arg Gln Gln Thr Asn
                                                               80
                                           75
                      70
  65
 Lys Ser Ile Gly Thr Tyr Gly Val Asn Tyr Pro Ala Asn Gly Asp Phe
                                                           95
                                       90
                  85
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800

Leu Ala Ala Ala Asp Gly Ala Asn Asp Ala Ser Asp His Ile Gln Gln

100 105 110

Met Ala Ser Ala Cys Arg Ala Thr Arg Leu Val Leu Gly Gly Tyr Ser 115 120 125

Gln Gly Ala Ala Val Ile Asp Ile Val Thr Ala Ala Pro Leu Pro Gly
130 135 140

Leu Gly Phe Thr Gln Pro Leu Pro Pro Ala Ala Asp Asp His Ile Ala 145 150 155 160

Ala Ile Ala Leu Phe Gly Asn Pro Ser Gly Arg Ala Gly Gly Leu Met 165 170 175

Ser Ala Leu Thr Pro Gln Phe Gly Ser Lys Thr Ile Asn Leu Cys Asn 180 185 190

Asn Gly Asp Pro Ile Cys Ser Asp Gly Asn Arg Trp Arg Ala His Leu 195 200 205

Gly Tyr Val Pro Gly Met Thr Asn Gln Ala Ala Arg Phe Val Ala Ser 210 215 220

Arg Ile 225

<210> 53

<211> 700

<212> DNA

<213> Mycobacterium tuberculosis

<400> 53

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<210> 54

<211> 181

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<212> PRT
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<213> Mycobacterium tuberculosis

<400> 54

Met Thr Glu Tyr Glu Gly Pro Lys Thr Lys Phe His Ala Leu Met Gln
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Glu Gln Ile His Asn Glu Phe Thr Ala Ala Gln Gln Tyr Val Ala Ile 20 25 30

Ala Val Tyr Phe Asp Ser Glu Asp Leu Pro Gln Leu Ala Lys His Phe 35 40 45

Tyr Ser Gln Ala Val Glu Glu Arg Asn His Ala Met Met Leu Val Gln 50 55 60

His Leu Leu Asp Arg Asp Leu Arg Val Glu Ile Pro Gly Val Asp Thr
65 70 75 80

Val Arg Asn Gln Phe Asp Arg Pro Arg Glu Ala Leu Ala Leu 85 90 95

Asp Gln Glu Arg Thr Val Thr Asp Gln Val Gly Arg Leu Thr Ala Val
100 105 110

Ala Arg Asp Glu Gly Asp Phe Leu Gly Glu Gln Phe Met Gln Trp Phe 115 120 125

Leu Gln Glu Gln Ile Glu Glu Val Ala Leu Met Ala Thr Leu Val Arg 130 135 140

Val Ala Asp Arg Ala Gly Ala Asn Leu Phe Glu Leu Glu Asn Phe Val 145 150 155 160

Ala Arg Glu Val Asp Val Ala Pro Ala Ala Ser Gly Ala Pro His Ala 165 170 175

Ala Gly Gly Arg Leu 180

<210> 55

<211> 950

<212> DNA

<213> Mycobacterium tuberculosis

<400> 55

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taaaattccq tcqtqaacaa tcqacccatc cqcctqctqa catccqqcaq qqctqqtttq 180
qqtqqqqqq cattqatcac cqccqtcqtc ctqctcatcq ccttqqqcqc tqtttqqacc 240
coggttgcct tcqccqatqq atqcccqqac qccqaaqtca cqttcqcccq cqqcaccqqc 300
gaqccqcccq qaatcgggcg cgttggccag gcgttcgtcg actcgctgcg ccagcagact 360
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qqaqacqqcq ccaacqacqc catatcqcac attaaqtcca tqqcctcqtc atqcccqaac 480
accaaqctqq tcttqqqqqq ctattcqcaq qqcqcaaccq tqatcqatat cqtqqcqqq 540
gttccgttgg gcagcatcag ctttggcagt ccgctacctg cggcatacgc agacaacgtc 600
qcaqcqqtcq cqqtcttcqq caatccqtcc aaccqcqccq qcqqatcqct qtcqaqcctq 660
agcccgctat teggttecaa ggcgattgac etgtgcaate ecacegatee gatetgccat 720
gtgggccccg gcaacgaatt cagcggacac atcgacggct acatacccac ctacaccacc 780
caggeggeta gtttegtegt geagaggete egegeegggt eggtgeeaca tetgeetgga 840
teegteeege agetgeeegg gtetgteett cagatgeeeg geaetgeege aeeggeteee 900
gaatcgctgc acggtcgctg acgctttgtc agtaagccca taaaatcgcg
                                                                  950
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<210> 56

<211> 262

<212> PRT

<213> Mycobacterium tuberculosis

<400> 56

Met Asn Asn Arg Pro Ile Arg Leu Leu Thr Ser Gly Arg Ala Gly Leu

1 5 10 15

Gly Ala Gly Ala Leu Ile Thr Ala Val Val Leu Leu Ile Ala Leu Gly
20 25 30

Ala Val Trp Thr Pro Val Ala Phe Ala Asp Gly Cys Pro Asp Ala Glu 35 40 45

Val Thr Phe Ala Arg Gly Thr Gly Glu Pro Pro Gly Ile Gly Arg Val
50 55 60

Gly Gln Ala Phe Val Asp Ser Leu Arg Gln Gln Thr Gly Met Glu Ile
65 70 75 80

Gly Val Tyr Pro Val Asn Tyr Ala Ala Ser Arg Leu Gln Leu His Gly
85 90 95

Gly Asp Gly Ala Asn Asp Ala Ile Ser His Ile Lys Ser Met Ala Ser 100 105 110

Ser Cys Pro Asn Thr Lys Leu Val Leu Gly Gly Tyr Ser Gln Gly Ala 115 120 125

Thr Val Ile Asp Ile Val Ala Gly Val Pro Leu Gly Ser Ile Ser Phe 130 135 140

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Gly Ser Pro Leu Pro Ala Ala Tyr Ala Asp Asn Val Ala Ala Val Ala
                                        155
                                                             160
145
                    150
Val Phe Gly Asn Pro Ser Asn Arg Ala Gly Gly Ser Leu Ser Ser Leu
                165
                                    170
Ser Pro Leu Phe Gly Ser Lys Ala Ile Asp Leu Cys Asn Pro Thr Asp
                                                     190
            180
                                185
Pro Ile Cys His Val Gly Pro Gly Asn Glu Phe Ser Gly His Ile Asp
                            200
        195
Gly Tyr Ile Pro Thr Tyr Thr Thr Gln Ala Ala Ser Phe Val Val Gln
    210
                        215
                                             220
Arg Leu Arg Ala Gly Ser Val Pro His Leu Pro Gly Ser Val Pro Gln
                    230
                                        235
                                                             240
Leu Pro Gly Ser Val Leu Gln Met Pro Gly Thr Ala Ala Pro Ala Pro
                                    250
                245
                                                         255
Glu Ser Leu His Gly Arg
            260
<210> 57
<211> 1000
<212> DNA
<213> Mycobacterium tuberculosis
<400> 57
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tocattaatt cactetetgg aacacceget gtagacetat ettettteae tgaetteetg 180
cgccgccagg cgccggagtt gctgccggca agcatcagcg gcggtgcgcc actcgcaggc 240
ggcgatgcgc aactgccgca cggcaccacc attgtcgcgc tgaaataccc cggcggtgtt 300
qtcatggcqg qtqaccgqcg ttcgacqcaq gqcaacatqa tttctqggcg tgatgtgcgc 360
aaggtgtata tcaccgatga ctacaccgct accggcatcg ctggcacggc tgcggtcgcg 420
gttgagtttg cccggctqta tgccgtggaa cttqaqcact acgagaagct cgagggtgtg 480
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gaccegcaga gegegggteg tategttteg ttegacgeeg ceggeggttg gaacategag 660
gaagaggget atcaggeggt gggetegggt tegetgtteg egaagtegte gatgaagaag 720
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<210> 58

<211> 291

<212> PRT

<213> Mycobacterium tuberculosis

<400> 58

Met Thr Trp Pro Leu Pro Asp Arg Leu Ser Ile Asn Ser Leu Ser Gly
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Thr Pro Ala Val Asp Leu Ser Ser Phe Thr Asp Phe Leu Arg Arg Gln 20 25 30

Ala Pro Glu Leu Leu Pro Ala Ser Ile Ser Gly Gly Ala Pro Leu Ala 35 40 45

Gly Gly Asp Ala Gln Leu Pro His Gly Thr Thr Ile Val Ala Leu Lys
50 55 60

Tyr Pro Gly Gly Val Val Met Ala Gly Asp Arg Arg Ser Thr Gln Gly 65 70 75 80

Asn Met Ile Ser Gly Arg Asp Val Arg Lys Val Tyr Ile Thr Asp Asp 85 90 95

Tyr Thr Ala Thr Gly Ile Ala Gly Thr Ala Ala Val Ala Val Glu Phe 100 105 110

Ala Arg Leu Tyr Ala Val Glu Leu Glu His Tyr Glu Lys Leu Glu Gly
115 120 125

Val Pro Leu Thr Phe Ala Gly Lys Ile Asn Arg Leu Ala Ile Met Val 130 135 140

Arg Gly Asn Leu Ala Ala Ala Met Gln Gly Leu Leu Ala Leu Pro Leu 145 150 155 160

Leu Ala Gly Tyr Asp Ile His Ala Ser Asp Pro Gln Ser Ala Gly Arg 165 170 175

Ile Val Ser Phe Asp Ala Ala Gly Gly Trp Asn Ile Glu Glu Gly
180 185 190

Tyr Gln Ala Val Gly Ser Gly Ser Leu Phe Ala Lys Ser Ser Met Lys 195 200 205

Lys Leu Tyr Ser Gln Val Thr Asp Gly Asp Ser Gly Leu Arg Val Ala

210 215 220

Val Glu Ala Leu Tyr Asp Ala Ala Asp Asp Ser Ala Thr Gly Gly
225 230 235 240

Pro Asp Leu Val Arg Gly Ile Phe Pro Thr Ala Val Ile Ile Asp Ala 245 250 255

Asp Gly Ala Val Asp Val Pro Glu Ser Arg Ile Ala Glu Leu Ala Arg 260 265 270

Ala Ile Ile Glu Ser Arg Ser Gly Ala Asp Thr Phe Gly Ser Asp Gly 275 280 285

Gly Glu Lys 290

<210> 59

<211> 900

<212> DNA

<213> Mycobacterium tuberculosis

<400> 59

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<210> 60

<211> 248

<212> PRT

<213> Mycobacterium tuberculosis

<400> 60

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1 5 10 15

Ser	Glu	Leu	Ala	Arg	Lys	Gly	Ile	Ala	Arg	Ala	Lys	Ser	Val	Val	Ala
			20					25					30		

Leu Ala Tyr Ala Gly Gly Val Leu Phe Val Ala Glu Asn Pro Ser Arg 35 40 45

Ser Leu Gln Lys Ile Ser Glu Leu Tyr Asp Arg Val Gly Phe Ala Ala 50 55 60

Ala Gly Lys Phe Asn Glu Phe Asp Asn Leu Arg Arg Gly Gly Ile Gln 65 70 75 80

Phe Ala Asp Thr Arg Gly Tyr Ala Tyr Asp Arg Arg Asp Val Thr Gly 85 90 95

Arg Gln Leu Ala Asn Val Tyr Ala Gln Thr Leu Gly Thr Ile Phe Thr 100 105 110

Glu Gln Ala Lys Pro Tyr Glu Val Glu Leu Cys Val Ala Glu Val Ala 115 120 125

His Tyr Gly Glu Thr Lys Arg Pro Glu Leu Tyr Arg Ile Thr Tyr Asp 130 135 140

Gly Ser Ile Ala Asp Glu Pro His Phe Val Val Met Gly Gly Thr Thr 145 150 155 160

Glu Pro Ile Ala Asn Ala Leu Lys Glu Ser Tyr Ala Glu Asn Ala Ser 165 170 175

Leu Thr Asp Ala Leu Arg Ile Ala Val Ala Ala Leu Arg Ala Gly Ser 180 185 190

Ala Asp Thr Ser Gly Gly Asp Gln Pro Thr Leu Gly Val Ala Ser Leu 195 200 205

Glu Val Ala Val Leu Asp Ala Asn Arg Pro Arg Arg Ala Phe Arg Arg 210 215 220

Ile Thr Gly Ser Ala Leu Gln Ala Leu Leu Val Asp Gln Glu Ser Pro 225 230 235 240

Gln Ser Asp Gly Glu Ser Ser Gly 245

<210> 61

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<211> 1560
<212> DNA
<213> Mycobacterium tuberculosis
<400> 61
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teteggagee ggteeeggeg ggtatgtege ggegattege geegeaeage teggeetgag 180
cactgcaatc gtcgaaccca agtactgggg cggagtatgc ctcaatgtcg gctgtatccc 240
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atttggcatc agcggcgagg tgaccttcga ctacggcatc gcctatgacc gcagccgaaa 360
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gatccacqqq tacqqcacat ttqccqacqc caacacqttq ttqqttqatc tcaacqacqq 480
eggtacagaa teggteaegt tegacaaege cateategeg aceggeagta geaceegget 540
qqttcccqqc acctcactqt cggccaacgt agtcacctac gaggaacaga tcctgtcccg 600
agagetgeeg aaategatea ttattgeegg agetggtgee attggeatgg agtteggeta 660
cgtgctgaag aactacggcg ttgacgtgac catcgtggaa ttccttccgc gggcgctgcc 720
caacgaggac gccgatgtgt ccaaggagat cgagaagcag ttcaaaaaagc tgggtgtcac 780
gatectgace gecaegaagg tegagteeat egeegatgge gggtegeagg teacegtgae 840
cgtcaccaag gacggcgtgg cgcaagagct taaggcggaa aaggtgttgc aggccatcgg 900
atttgcgccc aacgtcgaag ggtacgggct ggacaaggca ggcgtcgcgc tgaccgaccg 960
caaqqctatc qqtqtcqacq actacatqcq taccaacqtq qqccacatct acqctatcqg 1020
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cgaaaccatt gceggtgcag agactttgac gctgggcgac catcggatgt tgccgcgcgc 1140
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aggttacgac gtggtggtgg ccaagttccc gttcacggcc aacgccaagg cgcacggcgt 1260
gggtgacccc agtgggttcg tcaagctggt ggccgacgcc aagcacggcg agctactggg 1320
tgggcacctg gtcggccacg acgtggccga gctgctgccg gagctcacgc tggcgcagag 1380
gtgggacctg accgccagcg agctggctcg caacgtccac acccacccaa cgatgtctga 1440
ggcgctgcag gagtgcttcc acggcctggt tggccacatg atcaatttct gagcggctca 1500
tgacgaggeg cgcgagcact gacaccccc agatcatcat gggtgccatc ggtggtgtgg 1560
<210> 62
<211> 464
<212> PRT
<213> Mycobacterium tuberculosis
<400> 62
Met Thr His Tyr Asp Val Val Val Leu Gly Ala Gly Pro Gly Gly Tyr
Val Ala Ala Ile Arg Ala Ala Gln Leu Gly Leu Ser Thr Ala Ile Val
             20
                                 25
Glu Pro Lys Tyr Trp Gly Gly Val Cys Leu Asn Val Gly Cys Ile Pro
                             40
```

Ser Lys Ala Leu Leu Arg Asn Ala Glu Leu Val His Ile Phe Thr Lys

50 55 60

Asp	Ala	Lys	Ala	Phe	Gly	He	Ser	GLY	Glu	Val	Thr	Phe	Asp	Tyr	GIY
65					70					75					80

- Ile Ala Tyr Asp Arg Ser Arg Lys Val Ala Glu Gly Arg Val Ala Gly
 85 90 95
- Val His Phe Leu Met Lys Lys Asn Lys Ile Thr Glu Ile His Gly Tyr

 100 105 110
- Gly Thr Phe Ala Asp Ala Asn Thr Leu Leu Val Asp Leu Asn Asp Gly
 115 120 125
- Gly Thr Glu Ser Val Thr Phe Asp Asn Ala Ile Ile Ala Thr Gly Ser 130 135 140
- Ser Thr Arg Leu Val Pro Gly Thr Ser Leu Ser Ala Asn Val Val Thr 145 150 155 160
- Tyr Glu Glu Gln Ile Leu Ser Arg Glu Leu Pro Lys Ser Ile Ile Ile 165 170 175
- Ala Gly Ala Gly Ala Ile Gly Met Glu Phe Gly Tyr Val Leu Lys Asn 180 185 190
- Tyr Gly Val Asp Val Thr Ile Val Glu Phe Leu Pro Arg Ala Leu Pro 195 200 205
- Asn Glu Asp Ala Asp Val Ser Lys Glu Ile Glu Lys Gln Phe Lys Lys 210 215 220
- Leu Gly Val Thr Ile Leu Thr Ala Thr Lys Val Glu Ser Ile Ala Asp 225 230 235 240
- Gly Gly Ser Gln Val Thr Val Thr Val Thr Lys Asp Gly Val Ala Gln 245 250 255
- Glu Leu Lys Ala Glu Lys Val Leu Gln Ala Ile Gly Phe Ala Pro Asn 260 265 270
- Val Glu Gly Tyr Gly Leu Asp Lys Ala Gly Val Ala Leu Thr Asp Arg 275 280 285
- Lys Ala Ile Gly Val Asp Asp Tyr Met Arg Thr Asn Val Gly His Ile 290 295 300
- Tyr Ala Ile Gly Asp Val Asn Gly Leu Leu Gln Leu Ala His Val Ala

Glu Ala Gln Gly Val Val Ala Ala Glu Thr Ile Ala Gly Ala Glu Thr
325 330 335

Leu Thr Leu Gly Asp His Arg Met Leu Pro Arg Ala Thr Phe Cys Gln 340 345 350

Pro Asn Val Ala Ser Phe Gly Leu Thr Glu Gln Gln Ala Arg Asn Glu 355 360 365

Gly Tyr Asp Val Val Val Ala Lys Phe Pro Phe Thr Ala Asn Ala Lys 370 375 380

Ala His Gly Val Gly Asp Pro Ser Gly Phe Val Lys Leu Val Ala Asp 385 390 395 400

Ala Lys His Gly Glu Leu Leu Gly Gly His Leu Val Gly His Asp Val
405 410 415

Ala Glu Leu Pro Glu Leu Thr Leu Ala Gln Arg Trp Asp Leu Thr
420 425 430

Ala Ser Glu Leu Ala Arg Asn Val His Thr His Pro Thr Met Ser Glu
435 440 445

Ala Leu Gln Glu Cys Phe His Gly Leu Val Gly His Met Ile Asn Phe 450 455 460

<210> 63

<211> 550

<212> DNA

<213> Mycobacterium tuberculosis

<400> 63

ggeceggete geggecgee tgeaggaaaa gaaggeetge ceaggeceag acteageega 60 gtagteacee agtaceecae accaggaagg accgeccate atggeaaage tetecacega 120 cgaactgetg gaegegttea aggaaatgae cetgttggag etetecgaet tegteaagaa 180 gttegaggag acettegagg teacegeege eggeteeagte geegtegeeg eegeeggtge 240 cgeeceggee ggggaeaaga agateggegt eateaaggtg gteeggaga tegttteegg 360 cetgggeete aaggaegea aggaeetggt egaeggegg eecaageege tgetggagaa 420 ggtegeeaag gaggeegee aegaggeeaa ggeeaagetg gaggeegeeg gegecacegt 480 caeegteaag tagetetgee eagegtgtte ttttgegtet geteggeeeg tagegaacae 540

tgcgcccgct 550

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<211> 130

<212> PRT

<213> Mycobacterium tuberculosis

<400> 64

Met Ala Lys Leu Ser Thr Asp Glu Leu Leu Asp Ala Phe Lys Glu Met

1 5 10 15

Thr Leu Leu Glu Leu Ser Asp Phe Val Lys Lys Phe Glu Glu Thr Phe
20 25 30

Glu Val Thr Ala Ala Ala Pro Val Ala Val Ala Ala Ala Gly Ala Ala 35 40 45

Pro Ala Gly Ala Ala Val Glu Ala Ala Glu Glu Glu Ser Glu Phe Asp
50 55 60

Val Ile Leu Glu Ala Ala Gly Asp Lys Lys Ile Gly Val Ile Lys Val 65 70 75 80

Val Arg Glu Ile Val Ser Gly Leu Gly Leu Lys Glu Ala Lys Asp Leu 85 90 95

Val Asp Gly Ala Pro Lys Pro Leu Leu Glu Lys Val Ala Lys Glu Ala 100 105 110

Ala Asp Glu Ala Lys Ala Lys Leu Glu Ala Ala Gly Ala Thr Val Thr
115 120 125

Val Lys 130

<210> 65

<211> 900

<212> DNA

<213> Mycobacterium tuberculosis

<400> 65

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ttaacacgct gcaggtgccg gccggcatcg tccaggtgag ctaccacggt gatctcaccg 420 ccatcagegc ccgctcggaa tgggcacccg agttcgccat ccacgacctg gattcacttg 480 atgcgcttgc cgccgcac cccgccgact ttccggacga catcgcgcac tacctctgga 540 cctggaccga ccgctccgct ggctcgctgc gcgcccgcat gtttgccgcc aacttgggcg 600 tcaccgaaga cgaagcgacc ggtgccgcgg ccatccggat taccgattac ctcagccgtg 660 acctcaccat cacccagggc aaaggatcgt tgatccacac cacctggagt cccgagggct 720 gggttcgggt agccggcag gttgtcagcg acggtgtgc acaactcgac tgacgtagag 780 ctcagcgctg ccgatgcaac acggcggcaa ggtgatcctg caggggttgc ccgaccgcg 840 gcatctgcaa cgagtacgaa agctcgtcgc cgtcgatgcg gtaggaacgg tcaagggcgg 900

<210> 66

<211> 228

<212> PRT

<213> Mycobacterium tuberculosis

<400> 66

Met Ala Ile Glu Val Ser Val Leu Arg Val Phe Thr Asp Ser Asp Gly
1 5 10 15

Asn Phe Gly Asn Pro Leu Gly Val Ile Asn Ala Ser Lys Val Glu His
20 25 30

Arg Asp Arg Gln Gln Leu Ala Ala Gln Ser Gly Tyr Ser Glu Thr Ile 35 40 45

Phe Val Asp Leu Pro Ser Pro Gly Ser Thr Thr Ala His Ala Thr Ile 50 55 60

His Thr Pro Arg Thr Glu Ile Pro Phe Ala Gly His Pro Thr Val Gly 65 70 75 80

Ala Ser Trp Trp Leu Arg Glu Arg Gly Thr Pro Ile Asn Thr Leu Gln 85 90 95

Val Pro Ala Gly Ile Val Gln Val Ser Tyr His Gly Asp Leu Thr Ala 100 105 110

Ile Ser Ala Arg Ser Glu Trp Ala Pro Glu Phe Ala Ile His Asp Leu 115 120 125

Asp Ser Leu Asp Ala Leu Ala Ala Ala Asp Pro Ala Asp Phe Pro Asp 130 135 140

Asp Ile Ala His Tyr Leu Trp Thr Trp Thr Asp Arg Ser Ala Gly Ser 145 150 155 160

Leu Arg Ala Arg Met Phe Ala Ala Asn Leu Gly Val Thr Glu Asp Glu 165 170 175

50

```
Ala Thr Gly Ala Ala Ala Ile Arg Ile Thr Asp Tyr Leu Ser Arg Asp
Leu Thr Ile Thr Gln Gly Lys Gly Ser Leu Ile His Thr Trp Ser
                            200
                                                205
Pro Glu Gly Trp Val Arg Val Ala Gly Arg Val Val Ser Asp Gly Val
                        215
                                            220
Ala Gln Leu Asp
225
<210> 67
<211> 500
<212> DNA
<213> Mycobacterium tuberculosis
<400> 67
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ccggcgatgq qcatcggagg tgtggqtggt ttgggtgggg ccggttcggg tccggcgatg 120
ggcatgggg gtgtgggtgg tttgggtggg gccggttcgg gtccggcgat gggcatgggg 180
ggtgtgggtg gtttagatgc ggccggttcc ggcgagggcg gctctcctgc ggcgatcggc 240
atcggagttg gcggaggcgg aggtgggggt gggggtggcg gcggcggggc cgacacgaac 300
cgctccgaca ggtcgtcgga cgtcgggggc ggagtctggc cgttgggctt cggtaggttt 360
gccgatgcgg gcgccggcgg aaacgaagca ctggggtcga agaacggctg cgctgccata 420
tegteeggag ettecatace ttegtgegge eggaagaget tgtegtagte ggeegceatg 480
acaacctctc agagtgcgct
                                                                  500
<210> 68
<211> 139
<212> PRT
<213> Mycobacterium tuberculosis
<400> 68
Met Gly Ala Gly Pro Ala Met Gly Ile Gly Gly Val Gly Gly Leu Gly
                  5
                                                         15
Gly Ala Gly Ser Gly Pro Ala Met Gly Met Gly Val Gly Gly Leu
             20
                                 25
                                                     30
Gly Gly Ala Gly Ser Gly Pro Ala Met Gly Met Gly Gly Val Gly Gly
         35
                                                 45
                             40
Leu Asp Ala Ala Gly Ser Gly Glu Gly Ser Pro Ala Ala Ile Gly
```

60

55

```
65
                                              75
                           70
                                                                  80
      Ala Asp Thr Asn Arg Ser Asp Arg Ser Ser Asp Val Gly Gly Val
#
                                          90
                                                              95
                       85
      Trp Pro Leu Gly Phe Gly Arg Phe Ala Asp Ala Gly Ala Gly Gly Asn
                  100
                                      105
                                                         110
      Glu Ala Leu Gly Ser Lys Asn Gly Cys Ala Ala Ile Ser Ser Gly Ala
              115
                                  120
                                                     125
      Ser Ile Pro Ser Cys Gly Arg Lys Ser Leu Ser
          130
                              135
      <210> 69
      <211> 2050
      <212> DNA
      <213> Mycobacterium tuberculosis
      <400> 69
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      qqtatqqaaq ctccqqacqa tatqqcaqcq cagccqttct tcgaccccag tgcttcgttt 120
      cegceggege cegeategge aaacetaceg aageceaacg gecagaetee geeceegaeg 180
      teegaegaee tgteggageg gttegtgteg geeeggeege egeeaeeeee acceeeaeet 240
      cegecteege caacteegat geegategee geaggagage egeectegee ggaaceggee 300
      gcatetaaac cacccacacc ccccatgccc atcgccggac ccgaaccggc cccacccaaa 360
      ccacccacac cccccatgcc catcgccgga cccgaaccgg ccccacccaa accacccaca 420
      ceteogatge ceategeegg acetgeacce aceceaaccg aateceagtt ggegeecce 480
      agaccacega caccacaaac gccaacegga gcgccgcagc aaceggaatc aceggcgccc 540
      cacqtaccet cgcacqggcc acatcaacce cggcgcaccg caccagcacc gccctgggca 600
      aagatgecaa teggegaace eeegeeeget eegteeagae egtetgegte eeeggeegaa 660
      ccaccgaccc ggcctgccc ccaacactcc cgacgtgcgc gccggggtca ccgctatcgc 720
      acagacaceg aacgaaacgt cgggaaggta gcaactggtc catccatcca ggcgcggctg 780
      egggeagagg aageateegg egegeagete geeceeggaa eggageeete geeagegeeg 840
      ttgggccaac cgagategta tetggeteeg eecaceegee eegegeegae agaaceteee 900
      cccagcccct cgccgcagcg caactccggt cggcgtgccg agcgacgcgt ccaccccgat 960
      ttagccqccc aacatqccqc qqcqcaacct gattcaatta cggccqcaac cactqgcqgt 1020
      cgtcgccgca agcgtgcagc gccggatctc gacgcgacac agaaatcctt aaggccggcg 1080
      gccaaggggc cgaaggtgaa gaaggtgaag ccccagaaac cgaaggccac gaagccgccc 1140
      aaagtggtgt cgcagcgcgg ctggcgacat tgggtgcatg cgttgacgcg aatcaacctg 1200
      ggcctgtcac ccgacgagaa gtacgagetg gacctgcacg ctcgagtccg ccgcaatccc 1260
      cgcgggtcgt atcagatcgc cgtcgtcggt ctcaaaggtg gggctggcaa aaccacgctg 1320
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      geggatecag gegeeggaaa cetegeegat egggtaggge gacaateggg egegaceate 1440
      gctgatgtgc ttgcagaaaa agagctgtcg cactacaacg acatccgcgc acacactagc 1500
```

gtcaatgcgg tcaatctgqa agtgctqccg gcaccqqaat acagctcggc gcagcgcqcq 1560

Ser His Gly Pro His Gln Pro Arg Arg Thr Ala Pro Ala Pro Pro Trp
180 185 190

Ala Lys Met Pro Ile Gly Glu Pro Pro Pro Ala Pro Ser Arg Pro Ser 195 200 205

Ala Ser Pro Ala Glu Pro Pro Thr Arg Pro Ala Pro Gln His Ser Arg 210 215 220

Arg Ala Arg Arg Gly His Arg Tyr Arg Thr Asp Thr Glu Arg Asn Val 225 230 235 240

Gly Lys Val Ala Thr Gly Pro Ser Ile Gln Ala Arg Leu Arg Ala Glu 245 250 255

Glu Ala Ser Gly Ala Gln Leu Ala Pro Gly Thr Glu Pro Ser Pro Ala 260 265 270

Pro Leu Gly Gln Pro Arg Ser Tyr Leu Ala Pro Pro Thr Arg Pro Ala 275 280 285

Pro Thr Glu Pro Pro Pro Ser Pro Ser Pro Gln Arg Asn Ser Gly Arg 290 295 300

Arg Ala Glu Arg Arg Val His Pro Asp Leu Ala Ala Gln His Ala Ala 305 310 315 320

Ala Gln Pro Asp Ser Ile Thr Ala Ala Thr Thr Gly Gly Arg Arg Arg 325 330 335

Lys Arg Ala Ala Pro Asp Leu Asp Ala Thr Gln Lys Ser Leu Arg Pro 340 345 350

Ala Ala Lys Gly Pro Lys Val Lys Lys Val Lys Pro Gln Lys Pro Lys 355 360 365

Ala Thr Lys Pro Pro Lys Val Val Ser Gln Arg Gly Trp Arg His Trp 370 375 380

Val His Ala Leu Thr Arg Ile Asn Leu Gly Leu Ser Pro Asp Glu Lys 385 390 395 400

Tyr Glu Leu Asp Leu His Ala Arg Val Arg Arg Asn Pro Arg Gly Ser
405 410 415

Tyr Gln Ile Ala Val Val Gly Leu Lys Gly Gly Ala Gly Lys Thr Thr 420 425 430

Leu Thr Ala Ala Leu Gly Ser Thr Leu Ala Gln Val Arg Ala Asp Arg
435
440
445

Ile Leu Ala Leu Asp Ala Asp Pro Gly Ala Gly Asn Leu Ala Asp Arg 450 455 460

Val Gly Arg Gln Ser Gly Ala Thr Ile Ala Asp Val Leu Ala Glu Lys 465 470 475 480

Glu Leu Ser His Tyr Asn Asp Ile Arg Ala His Thr Ser Val Asn Ala 485 490 495

Val Asn Leu Glu Val Leu Pro Ala Pro Glu Tyr Ser Ser Ala Gln Arg
500 505 510

Ala Leu Ser Asp Ala Asp Trp His Phe Ile Ala Asp Pro Ala Ser Arg
515 520 525

Phe Tyr Asn Leu Val Leu Ala Asp Cys Gly Ala Gly Phe Phe Asp Pro 530 535 540

Leu Thr Arg Gly Val Leu Ser Thr Val Ser Gly Val Val Val Ala 545 550 555 560

Ser Val Ser Ile Asp Gly Ala Gln Gln Ala Ser Val Ala Leu Asp Trp
565 570 575

Leu Arg Asn Asn Gly Tyr Gln Asp Leu Ala Ser Arg Ala Cys Val Val 580 585 590

Ile Asn His Ile Met Pro Gly Glu Pro Asn Val Ala Val Lys Asp Leu 595 600 605

Val Arg His Phe Glu Gln Gln Val Gln Pro Gly Arg Val Val Met 610 620

Pro Trp Asp Arg His Ile Ala Ala Gly Thr Glu Ile Ser Leu Asp Leu 625 630 635 640

Leu Asp Pro Ile Tyr Lys Arg Lys Val Leu Glu Leu Ala Ala Ala Leu 645 650 655

Ser Asp Asp Phe Glu Arg Ala Gly Arg Arg 660 665

<210> 71

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<213> Mycobacterium tuberculosis
  <400> 71
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  ctggaccagc tcggcactgc tgaatcgcgt gcgtacaaga tgtggctgcc gccgttgacc 180
  aatccggtcc cgctcaacga gctcatcgcc cgtgatcggc gacaacccct gcgatttgcc 240
  ctggggatca tggatgaacc gcgccgccat ctacaggatg tgtggggcgt agacgtttcc 300
  ggggccggcg gcaacatcgg tattgggggc gcacctcaaa ccgggaagtc gacgctactg 360
  cagacgatgg tgatgtcggc cgccgccaca cactcaccgc gcaacgttca gttctattgc 420
  atcgacctag gtggcggcgg gctgatctat ctcgaaaacc ttccacacgt cggtggggta 480
  gccaatcggt ccgagcccga caaggtcaac cgggtggtcg cagagatgca agccgtcatg 540
  cggcaacggg aaaccacctt caaggaacac cgagtgggct cgatcgggat gtaccggcag 600
  ctgcgtgacg atccaagtca acccgttgcg tccgatccat acggcgacgt ctttctgatc 660
  atcgacggat ggcccggttt tgtcggcgag ttccccgacc ttgaggggca ggttcaagat 720
ctggccgccc aggggctggg gttcggcgtc cacgtcatca tctccacgcc acgctggaca 780
gagetgaagt egegtgtteg egactaeete ggeaceaaga tegagtteeg gettggtgae 840
  gtcaatgaaa cccagatcga ccggattacc cgcgagatcc cggcgaatcg tccgggtcgg 900
  gcagtgtcga tggaaaagca ccatctgatg atcggcgtgc ccaggttcga cggcgtgcac 960
  agegeegata acetggtgga ggegateace gegggggtga egeagatege tteecageae 1020
  accgaacagg cacctccggt gcgggtcctg ccggagcgta tccacctgca cgaactcgac 1080
ľŌ
  ccqaacccgc cgggaccaga gtccgactac cgcactcgct gggagattcc gatcggcttg 1140
   cgcgagacgg acctgacgcc ggctcactgc cacatgcaca cgaacccgca cctactgatc 1200
ttcggtgcgg ccaaatcggg caagacgacc attgcccacg cgatcgcgcg cgccatttgt 1260
gcccgaaaca gtccccagca ggtgcggttc atgctcgcgg actaccgctc gggcctgctg 1320
🛓 gacgeggtge eggacaceca tetgetggge geeggegega teaacegeaa eagegegteg 1380
ctagacgagg ccgctcaagc actggcggtc aacctgaaga agcggttgcc gccgaccgac 1440
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   gtcgacgatt ggcacatgat cgtgggtgcc gccgggggga tgccgccgat ggcaccgctg 1560
   gccccgttat tgccggcggc ggcagatatc gggttgcaca tcattgtcac ctgtcagatg 1620
   agccaggett acaaggcaac catggacaag ttegteggeg cegeattegg gtegggeget 1680
   ccgacaatgt tcctttcggg cgagaagcag gaattcccat ccagtgagtt caaggtcaag 1740
   cggcgccccc ctggccaggc atttctcgtc tcgccagacg gcaaagaggt catccaggcc 1800
   ccctacatcg agcctccaga agaagtgttc gcagcacccc caagcgccgg ttaagattat 1860
                                                                     1890
   ttcattgccg gtgtagcagg acccgagctc
   <210> 72
   <211> 591
   <212> PRT
   <213> Mycobacterium tuberculosis
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<211> 1890 <212> DNA

1

Gln Leu Gly Thr Ala Glu Ser Arg Ala Tyr Lys Met Trp Leu Pro Pro

Met Thr Ala Glu Pro Glu Val Arg Thr Leu Arg Glu Val Val Leu Asp

10

5

15

275 280 285

Arg	Phe 290	Asp	Gly	Val	His	Ser 295	Ala	Asp	Asn	Leu	Val 300	Glu	Ala	Ile	Thr
Ala 305	Gly	Val	Thr	Gln	Ile 310	Ala	Ser	Gln	His	Thr 315	Glu	Gln	Ala	Pro	Pro 320
Val	Arg	Val	Leu	Pro 325	Glu	Arg	Ile	His	Leu 330	His	Glu	Leu	Asp	Pro 335	Asn
Pro	Pro	Gly	Pro 340	Glu	Ser	Asp	Tyr	Arg 345	Thr	Arg	Trp	Glu	Ile 350	Pro	Ile
Gly	Leu	Arg 355	Glu	Thr	Asp	Leu	Thr 360	Pro	Ala	His	Суз	His 365	Met	His	Thr
Asn	Pro 370	His	Leu	Leu	Ile	Phe 375	Gly	Ala	Ala	Lys	Ser 380	Gly	Lys	Thr	Thr
Ile 385	Ala	His	Ala	Ile	Ala 390	Arg	Ala	Ile	Cys	Ala 395	Arg	Asn	Ser	Pro	Gln 400
Gln	Val	Arg	Phe	Met 405	Leu	Ala	Asp	Tyr	Arg 410	Ser	Gly	Leu	Leu	Asp 415	Ala
Val	Pro	Asp	Thr 420	His	Leu	Leu	Gly	Ala 425	Gly	Ala	Ile	Asn	Arg 430	Asn	Ser
Ala	Ser	Leu 435	Asp	Glu	Ala	Ala	Gln 440	Ala	Leu	Ala	Val	Asn 445	Leu	Lys	Lys
Arg	Leu 450	Pro	Pro	Thr	Asp	Leu 455	Thr	Thr	Ala	Gln	Leu 460	Arg	Ser	Arg	Ser
Trp 465	Trp	Ser	Gly	Phe	Asp 470	Val	Val	Leu	Leu	Val 475	Asp	Asp	Trp	His	Met 480
Ile	Val	Gly	Ala	Ala 485	Gly	Gly	Met	Pro	Pro 490	Met	Ala	Pro	Leu	Ala 495	Pro
Leu	Leu	Pro	Ala 500	Ala	Ala	Asp	Ile	Gly 505	Leu	His	Ile	Ile	Val 510	Thr	Cys
Gln	Met	Ser 515	Gln	Ala	Tyr	Lys	Ala 520	Thr	Met	Asp	Lys	Phe 525	Val	Gly	Ala

Ala Phe Gly Ser Gly Ala Pro Thr Met Phe Leu Ser Gly Glu Lys Gln

1.4

14

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Glu Phe Pro Ser Ser Glu Phe Lys Val Lys Arg Arg Pro Pro Gly Gln
                    550
                                         555
Ala Phe Leu Val Ser Pro Asp Gly Lys Glu Val Ile Gln Ala Pro Tyr
                565
                                     570
Ile Glu Pro Pro Glu Glu Val Phe Ala Ala Pro Pro Ser Ala Gly
            580
                                585
<210> 73
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<212> PRT
<213> Mycobacterium tuberculosis
<400> 73
Asp Pro Val Asp Asp Ala Phe Ile Ala Lys Leu Asn Thr Ala Gly
                  5
                                      10
<210> 74
<211> 14
<212> PRT
<213> Mycobacterium tuberculosis
<220>
<221> UNSURE
<222> (14)
<223> Xaa is unknown
<400> 74
Asp Pro Val Asp Ala Ile Ile Asn Leu Asp Asn Tyr Gly Xaa
                  5
                                      10
<210> 75
<211> 15
<212> PRT
<213> Mycobacterium tuberculosis
<220>
<221> UNSURE
<222> (5)
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<223> Xaa is unknown

<400> 75

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Ala Glu Met Lys Xaa Phe Lys Asn Ala Ile Val Gln Glu Ile Asp
                                      10
                   5
  1
<210> 76
<211> 14
<212> PRT
<213> Mycobacterium tuberculosis
<220>
<221> VARIANT
<222> (3)
<223> Ala is Ala or Gln
<220>
<221> VARIANT
<222> (7)
<223> Thr is Gly or Thr
<220>
<221> UNSURE
<222> (11)
<223> Xaa is unknown
<400> 76
Val Ile Ala Gly Met Val Thr His Ile His Xaa Val Ala Gly
                   5
<210> 77
 <211> 15
 <212> PRT
 <213> Mycobacterium tuberculosis
 <400> 77
 Thr Asn Ile Val Val Leu Ile Lys Gln Val Pro Asp Thr Trp Ser
                                                            15
                                       10
                   5
   1
 <210> 78
 <211> 15
 <212> PRT
 <213> Mycobacterium tuberculosis
 <400> 78
 Ala Ile Glu Val Ser Val Leu Arg Val Phe Thr Asp Ser Asp Gly
                                                            15
                                       10
                    5
```

<212> PRT

```
<210> 79
<211> 15
<212> PRT
<213> Mycobacterium tuberculosis
<400> 79
Ala Lys Leu Ser Thr Asp Glu Leu Leu Asp Ala Phe Lys Glu Met
                                      10
                  5
<210> 80
<211> 15
<212> PRT
<213> Mycobacterium tuberculosis
<220>
<221> VARIANT
<222> (4)
<223> Asp is Asp or Glu
<400> 80
Asp Pro Ala Asp Ala Pro Asp Val Pro Thr Ala Ala Gln Leu Thr
                                                           15
                                      10
  1
                   5
<210> 81
<211> 50
<212> PRT
<213> Mycobacterium tuberculosis
Ala Glu Asp Val Arg Ala Glu Ile Val Ala Ser Val Leu Glu Val Val
                                                           15
                                      10
                   5
  1
Val Asn Glu Gly Asp Gln Ile Asp Lys Gly Asp Val Val Leu Leu
                                  25
                                                       30
              20
Glu Ser Met Tyr Met Glu Ile Pro Val Leu Ala Glu Ala Ala Gly Thr
                                                   45
                              40
          35
 Val Ser
      50
 <210> 82
 <211> 15
```

```
The Half of the Control of the Last Control of the Control of the
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<213> Mycobacterium tuberculosis
<400> 82
Thr Thr Ser Pro Asp Pro Tyr Ala Ala Leu Pro Lys Leu Pro Ser
                                                           15
                                      10
                  5
<210> 83
<211> 15
<212> PRT
<213> Mycobacterium tuberculosis
<400> 83
Thr Glu Tyr Glu Gly Pro Lys Thr Lys Phe His Ala Leu Met Gln
                   5
<210> 84
<211> 15
<212> PRT
<213> Mycobacterium tuberculosis
<400> 84
Thr Thr Ile Val Ala Leu Lys Tyr Pro Gly Gly Val Val Met Ala
                                                           15
                                      10
                   5
<210> 85
<211> 15
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 <221> UNSURE
 <222> (10)
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acatgatccg atcgctgccg acattggcac gcaagtgagc gacaacgctc tgcacggcgt 180
gaeggeegge tegaeggege tgaegteggt gaeeggetg gtteeegegg gggeegatga 240
gqtctccgcc caagcggcga cggcgttcac atcggagggc atccaattgc tggcttccaa 300
tgcatcggcc caagaccagc tccaccgtgc gggcgaagcg gtccaggacg tcgcccgcac 360
ctattcgcaa atcgacgacg gcgccgccgg cgtcttcgcc taataggccc ccaacacatc 420
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ggagggagtg atcaccatgc tgtggcacgc
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<212> PRT
<213> Mycobacterium tuberculosis
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                                                      30
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Thr Ser Val Thr Gly Leu Val Pro Ala Gly Ala Asp Glu Val Ser Ala
                                                  45
                              40
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Gln Ala Ala Thr Ala Phe Thr Ser Glu Gly Ile Gln Leu Leu Ala Ser
                          55
     50
Asn Ala Ser Ala Gln Asp Gln Leu His Arg Ala Gly Glu Ala Val Gln
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                  85
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ageggatggt tegacagegg aetggtgeeg ageaggeeca tetgegegge tteetegteg 180
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gccgccggcg gcaactggcc aaccggtgtt gagctgccag gggagggcat tccgaagatc 360
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Asp Ser Gly Leu Val Pro Ser Arg Pro Ile Cys Ala Ala Ser Ser Ser
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Ala Gly Leu Pro Pro Pro Val Pro Pro Thr Trp Leu Asn Asn Asp Val
                                              60
                          55
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Thr Cys Cys Ser Gly Trp Val Ser Cys Cys Ile Gly Pro Leu Ile Ser
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                                                               80
                      70
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Pro Ser Trp Pro Arg Val Trp Val Ala Ala Gly Gly Asn Trp Pro Thr
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Gly Val Glu Leu Pro Gly Glu Gly Ile Pro Lys Ile Gly Phe Val Val
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120

115

125

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15

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Thr Ala Arg Leu Asn Ser Leu Gly Glu Ala Trp Thr Gly Gly Ser

Asp 65	Lys	Ala	Leu	Ala	Ala 70	Ala	Thr	Pro	Met	Val 75	Val	Trp	Leu	Gln	Thr 80
Ala	Ser	Thr	Gln	Ala 85	Lys	Thr	Arg	Ala	Met 90	Gln	Ala	Thr	Ala	Gln 95	Ala
Ala	Ala	Tyr	Thr 100	Gln	Ala	Met	Ala	Thr 105	Thr	Pro	Ser	Leu	Pro 110	Glu	Ile
Ala	Ala	Asn 115	His	Ile	Thr	Gln	Ala 120	Val	Leu	Thr	Ala	Thr 125	Asn	Phe	Phe
Gly	Ile 130	Asn	Thr	Ile	Pro	Ile 135	Ala	Leu	Thr	Glu	Met 140	Asp	Tyr	Phe	Ile
Arg 145	Met	Trp	Asn	Gln	Ala 150	Ala	Leu	Ala	Met	Glu 155	Val	Tyr	Gln	Ala	Glu 160
Thr	Ala	Val	Asn	Thr 165	Leu	Phe	Glu	Lys	Leu 170	Glu	Pro	Met	Ala	Ser 175	Ile
Leu	Asp	Pro	Gly 180	Ala	Ser	Gln	Ser	Thr 185	Thr	Asn	Pro	Ile	Phe 190	Gly	Met
Pro	Ser	Pro 195	Gly	Ser	Ser	Thr	Pro 200	Val	Gly	Gln	Leu	Pro 205	Pro	Ala	Ala
Thr	Gln 210	Thr	Leu	Gly	Gln	Leu 215	Gly	Glu	Met	Ser	Gly 220	Pro	Met	Gln	Gln
Leu 225		Gln	Pro	Leu	Gln 230		Val	Thr	Ser	Leu 235		Ser	Gln	Val	Gly 240
Gly	Thr	Gly	Gly	Gly 245		Pro	Ala	Asp	Glu 250	Glu	Ala	Ala	Gln	Met 255	Gly
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Gly	Pro	Ser 275		. Gly	Ala	Gly	Leu 280		Arg	Ala	Glu	Ser 285	Leu	Pro	Gly
Ala	Gly 290	_	Ser	Leu	Thr	Arg 295		Pro	Leu	Met	Ser 300		Leu	Ile	Glu

Lys Pro Val Ala Pro Ser Val Met Pro Ala Ala Ala Gly Ser Ser

tion from the first term of th

Ala Thr Gly Gly Ala Ala Pro Val Gly Ala Gly Ala Met Gly Gln Gly 325 330 335

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Asp Asp Trp 370

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- Gly Leu Lys Gly Gly Ala Gly Lys Thr Thr Leu Thr Ala Ala Leu Gly 65 70 75 80
- Ser Thr Leu Ala Gln Val Arg Ala Asp Arg Ile Leu Ala Leu Asp Ala 85 90 95
- Asp Pro Gly Ala Gly Asn Leu Ala Asp Arg Val Gly Arg Gln Ser Gly
 100 105 110
- Ala Thr Ile Ala Asp Val Leu Ala Glu Lys Glu Leu Ser His Tyr Asn 115 120 125
- Asp Ile Arg Ala His Thr Ser Val Asn Ala Val Asn Leu Glu Val Leu 130 135 140
- Trp His Phe Ile Ala Asp Pro Ala Ser Arg Phe Tyr Asn Leu Val Leu 165 170 175
- Ala Asp Cys Gly Ala Gly Phe Phe Asp Pro Leu Thr Arg Gly Val Leu 180 185 190
- Ser Thr Val Ser Gly Val Val Val Val Ala Ser Val Ser Ile Asp Gly
 195 200 205
- Ala Gln Gln Ala Ser Val Ala Leu Asp Trp Leu Arg Asn Asn Gly Tyr 210 215 220
- Gln Asp Leu Ala Ser Arg Ala Cys Val Val Ile Asn His Ile Met Pro 225 230 235 240
- Gly Glu Pro Asn Val Ala Val Lys Asp Leu Val Arg His Phe Glu Gln 245 250 255
- Gln Val Gln Pro Gly Arg Val Val Wet Pro Trp Asp Arg His Ile 260 265 270

Ala Al	a Gly 275	Thr	Glu	Ile	Ser	Leu 280	Asp	ьeu	ьeu	Asp	285	ше	ıyr	пуѕ	
Arg Ly 29		Leu	Glu	Leu	Ala 295	Ala	Ala	Leu	Ser	Asp 300	Asp	Phe	Glu	Arg	
Ala Gl 305	y Arg	Arg													
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Val Ser	
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gaagagtgtc tcggccagcg gcgatacctt gggtgccgtc atcagcgacc tggaggccaa 180
ctattcgggc atttccgagc gcctgatgga cccgtcttcc ccaggtaagt tgcaccgctt 240
cqtqaacatc tacqtcaacq acqaggacqt gcggttctcc ggcggcttgg ccaccgcgat 300
cgctgacggt gactcggtca ccatcctccc cgccgtggcc ggtgggtgag cggagcacat 360
gacacgatac gactegetgt tgcaggectt gggcaacacg ccgctggttg gcctgcagcg 420
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Met Asn Val Thr Val Ser Ile Pro Thr Ile Leu Arg Pro His Thr Gly
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Gly Gln Lys Ser Val Ser Ala Ser Gly Asp Thr Leu Gly Ala Val Ile
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Ser Asp Leu Glu Ala Asn Tyr Ser Gly Ile Ser Glu Arg Leu Met Asp
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Pro Ser Ser Pro Gly Lys Leu His Arg Phe Val Asn Ile Tyr Val Asn 50 55 60
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<211> 98

<212> PRT

<213> Mycobacterium tuberculosis

<400> 143

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Gly Arg Phe Glu Val His Ala Gln Thr Val Glu Asp Glu Ala Arg Arg 20 25 30

Met Trp Ala Ser Ala Gln Asn Ile Ser Gly Ala Gly Trp Ser Gly Met 35 40 45

Ala Glu Ala Thr Ser Leu Asp Thr Met Ala Gln Met Asn Gln Ala Phe
50 55 60

Arg Asn Ile Val Asn Met Leu His Gly Val Arg Asp Gly Leu Val Arg 65 70 75 80

Asp Ala Asn Asn Tyr Glu Gln Glu Gln Ala Ser Gln Gln Ile Leu

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85 90 95

Ser Ser

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<211> 940
<212> DNA
<213> Mycobacterium tuberculosis
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<400> 144 gccccagtc

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gccccagtcc tcgatcgcct catcgccttc accggccgcc agccgaccgc aggccacgtg 60
tecgecacet aacgaaagga tgateatgee caagagaage gaatacagge aaggeacgee 120
gaactgggtc gaccttcaga ccaccgatca gtccgccgcc aaaaagttct acacatcgtt 180
gtteggetgg ggttacgacg acaacceggt ccceggaggc ggtggggtct attccatggc 240
cacgetgaac ggcgaagecg tggccgccat cgcaccgatg cccccgggtg caccggaggg 300
gatgccqccq atctqgaaca cctatatcgc ggtggacgac gtcgatgcgg tggtggacaa 360
ggtggtgccc gggggcggcc aggtgatgat gccggccttc gacatcggcg atgccggccg 420
gatgtegtte atcacegate egaceggege tgeegtggge etatggeagg ceaateggea 480
categgageg aegttggtea aegagaeggg caegeteate tggaaegaae tgeteaegga 540
caageeggat ttggegetag egttetaega ggetgtggtt ggeeteaece actegageat 600
ggagataget gegggeeaga actategggt geteaaggee ggegaegegg aagteggegg 660
ctgtatggaa ccgccgatgc ccggcgtgcc gaatcattgg cacgtctact ttgcggtgga 720
tgacgccgac gccacggcgg ccaaagccgc cgcagcgggc ggccaggtca ttgcggaacc 780
ggctgacatt ccgtcggtgg gccggttcgc cgtgttgtcc gatccgcagg gcgcgatctt 840
cagtgtgttg aagcccgcac cgcagcaata gggagcatcc cgggcaggcc cgccggccgg 900
                                                                  940
cagattegga gaatgetaga agetgeegee ggegeegeeg
```

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<210> 145
<211> 261
<212> PRT
<213> Mycobacterium tuberculosis
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<400> 145

Met Pro Lys Arg Ser Glu Tyr Arg Gln Gly Thr Pro Asn Trp Val Asp
1 5 10 15

Leu Gln Thr Thr Asp Gln Ser Ala Ala Lys Lys Phe Tyr Thr Ser Leu 20 25 30

Phe Gly Trp Gly Tyr Asp Asp Asn Pro Val Pro Gly Gly Gly Val 35 40 45

Tyr Ser Met Ala Thr Leu Asn Gly Glu Ala Val Ala Ala Ile Ala Pro 50 55 60

Met Pro Pro Gly Ala Pro Glu Gly Met Pro Pro Ile Trp Asn Thr Tyr

THE STATE OF THE REAL PROPERTY.

47

Tige Special

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14

65	70	75	80

Ile Ala Val Asp Asp Val Asp Ala Val Val Asp Lys Val Val Pro Gly
85 90 95

Gly Gly Gln Val Met Met Pro Ala Phe Asp Ile Gly Asp Ala Gly Arg
100 105 110

Met Ser Phe Ile Thr Asp Pro Thr Gly Ala Ala Val Gly Leu Trp Gln
115 120 125

Ala Asn Arg His Ile Gly Ala Thr Leu Val Asn Glu Thr Gly Thr Leu 130 135 140

Tyr Glu Ala Val Val Gly Leu Thr His Ser Ser Met Glu Ile Ala Ala 165 170 175

Gly Gln Asn Tyr Arg Val Leu Lys Ala Gly Asp Ala Glu Val Gly 180 185 190

Cys Met Glu Pro Pro Met Pro Gly Val Pro Asn His Trp His Val Tyr 195 200 205

Phe Ala Val Asp Asp Ala Asp Ala Thr Ala Ala Lys Ala Ala Ala Ala 210 215 220

Gly Gly Gln Val Ile Ala Glu Pro Ala Asp Ile Pro Ser Val Gly Arg 225 230 235 240

Phe Ala Val Leu Ser Asp Pro Gln Gly Ala Ile Phe Ser Val Leu Lys 245 250 255

Pro Ala Pro Gln Gln 260

<210> 146

<211> 280

<212> DNA

<213> Mycobacterium tuberculosis

<400> 146

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<400> 149

<212> PRT

<213> Mycobacterium tuberculosis

Met Asn Leu Arg Arg His Gln Thr Leu Thr Leu Arg Leu Leu Ala Ala

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io
E.A.
122
#1 :t=:
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Series Series
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Ser Ala Gly Ile Leu Ser Ala Ala Ala Phe Ala Ala Pro Ala Gln Ala 20 25 30

Asn Pro Val Asp Asp Ala Phe Ile Ala Ala Leu Asn Asn Ala Gly Val
35 40 45

Asn Tyr Gly Asp Pro Val Asp Ala Lys Ala Leu Gly Gln Ser Val Cys
50 55 60

Pro Ile Leu Ala Glu Pro Gly Gly Ser Phe Asn Thr Ala Val Ala Ser 65 70 75 80

Val Val Ala Arg Ala Gln Gly Met Ser Gln Asp Met Ala Gln Thr Phe 85 90 95

Thr Ser Ile Ala Ile Ser Met Tyr Cys Pro Ser Val Met Ala Asp Val
100 105 110

Ala Ser Gly Asn Leu Pro Ala Leu Pro Asp Met Pro Gly Leu Pro Gly
115 120 125

Ser

<210> 150

<211> 400

<212> DNA

<213> Mycobacterium tuberculosis

<400> 150

atagtttggg gaaggtgtc ataaatgagg ctgtcgttga ccgcattgag cgccggtgta 60 ggcgccgtgg caatgtcgt gaccgtcggg gccggggtcg cctccgcaga tcccgtggac 120 gcggtcatta acaccactg caattacggg caggtagtag ctgcgctcaa cgcgacggat 180 ccgggggctg ccgcacagtt caacgcctca ccggtggcgc agtcctattt gcgcaatttc 240 ctcgccgcac cgccacctca gcgcgctgcc atggccgcgc aattgcaagc tgtgccgggg 300 gcggcacagt acatcggcct tgtcgagtcg gttgccggct cctgcaacaa ctattaagcc 360 catgcgggcc ccatcccgc acccggaac gtcgccggg

<210> 151

<211> 110

<212> PRT

<213> Mycobacterium tuberculosis

<400> 151

Met Arg Leu Ser Leu Thr Ala Leu Ser Ala Gly Val Gly Ala Val Ala

Met Ser Leu Thr Val Gly Ala Gly Val Ala Ser Ala Asp Pro Val Asp 20 25 30

Ala Val Ile Asn Thr Thr Cys Asn Tyr Gly Gln Val Val Ala Ala Leu 35 40 45

Asn Ala Thr Asp Pro Gly Ala Ala Ala Gln Phe Asn Ala Ser Pro Val
50 55 60

Ala Gln Ser Tyr Leu Arg Asn Phe Leu Ala Ala Pro Pro Pro Gln Arg
65 70 75 80

Ala Ala Met Ala Ala Gln Leu Gln Ala Val Pro Gly Ala Ala Gln Tyr 85 90 95

Ile Gly Leu Val Glu Ser Val Ala Gly Ser Cys Asn Asn Tyr
100 105 110

<210> 152

<211> 990

<212> DNA

<213> Mycobacterium tuberculosis

<400> 152

aataqtaata tcqctgtqcq gttgcaaaac gtgtgaccga ggttccgcag tcgagcgctg 60 cgggccgcct tcgaggagga cgaaccacag tcatgacgaa catcgtggtc ctgatcaagc 120 aggtcccaga tacctggtcg gagcgcaagc tgaccgacgg cgatttcacg ctggaccgcg 180 aggcegeega egeggtgetg gaegagatea aegagegege egtggaggaa gegetaeaga 240 ttcgggagaa agaggccgcc gacggcatcg aagggtcggt aaccgtgctg acggcgggcc 300 ccgaqcgcgc caccgaggcg atccgcaagg cgctgtcgat gggtgccgac aaggccgtcc 360 acctaaagga cgacggcatg cacggctcgg acgtcatcca aaccgggtgg gctttggcgc 420 gegegttggg caccategag ggeacegage tggtgatege aggeaaegaa tegacegaeg 480 gggtgggcgg tgcggtgccg gccatcatcg ccgagtacct gggcctgccg cagctcaccc 540 acctgcgcaa agtgtcgatc gagggcggca agatcaccgg cgagcgtgag accgatgagg 600 qcgtattcac cctcqaqqcc acqctqcccg cggtqatcag cgtgaacgag aagatcaacg 660 agecgegett eccepteette aaaggeatea tggeegeeaa gaagaaggaa gttacegtge 720 tqaccctqqc cqaqatcqqt qtcqaqaqcg acqaqqtqqq qctqgccaac gccggatcca 780 ccgtgctggc gtcgacgccc aaaccggcca agactgccgg ggagaaggtc accgacgagg 840 gtgaaggegg caaccagate gtgeagtace tggttgeeca gaaaatcate taagacatac 900 gcacctccca aagacgagag cgatataacc catggctgaa gtactggtgc tcgttgagca 960 990 cgctgaaggc gcgttaaaga aggtcagcgc

<210> 153

<211> 266

<212> PRT

<213> Mycobacterium tuberculosis

	-	_			_	_
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- Met Thr Asn Ile Val Val Leu Ile Lys Gln Val Pro Asp Thr Trp Ser

 1 5 10 15
- Glu Arg Lys Leu Thr Asp Gly Asp Phe Thr Leu Asp Arg Glu Ala Ala 20 25 30
- Asp Ala Val Leu Asp Glu Ile Asn Glu Arg Ala Val Glu Glu Ala Leu
 35 40 45
- Gln Ile Arg Glu Lys Glu Ala Ala Asp Gly Ile Glu Gly Ser Val Thr
 50 55 60
- Val Leu Thr Ala Gly Pro Glu Arg Ala Thr Glu Ala Ile Arg Lys Ala 65 70 75 80
- Leu Ser Met Gly Ala Asp Lys Ala Val His Leu Lys Asp Asp Gly Met 85 90 95
- His Gly Ser Asp Val Ile Gln Thr Gly Trp Ala Leu Ala Arg Ala Leu 100 105 110
- Gly Thr Ile Glu Gly Thr Glu Leu Val Ile Ala Gly Asn Glu Ser Thr 115 120 125
- Asp Gly Val Gly Gly Ala Val Pro Ala Ile Ile Ala Glu Tyr Leu Gly 130 135 140
- Leu Pro Gln Leu Thr His Leu Arg Lys Val Ser Ile Glu Gly Gly Lys
 145 150 155 160
- Ile Thr Gly Glu Arg Glu Thr Asp Glu Gly Val Phe Thr Leu Glu Ala 165 170 175
- Thr Leu Pro Ala Val Ile Ser Val Asn Glu Lys Ile Asn Glu Pro Arg 180 185 190
- Phe Pro Ser Phe Lys Gly Ile Met Ala Ala Lys Lys Glu Val Thr 195 200 205
- Val Leu Thr Leu Ala Glu Ile Gly Val Glu Ser Asp Glu Val Gly Leu 210 215 220
- Ala Asn Ala Gly Ser Thr Val Leu Ala Ser Thr Pro Lys Pro Ala Lys 225 230 235 240

260 265	
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<212> DNA	
<213> Mycobacterium tuberculosis	
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ctcccatggt accctaggac ccgggcagcc ccggc	35
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<211> 29	
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ctgagatcta tgaggctgtc gttgaccgc	29
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<211> 30	
<212> DNA	
<213> Mycobacterium tuberculosis	
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ctcccgggc ttaatagttg ttgcaggagc	30
<210> 158	
<211> 33	
<212> DNA	
<213> Mycobacterium tuberculosis	
<400> 158	
gcttagatct atgattttct gggcaaccag gta	33
·	
-210- 150	

Thr Ala Gly Glu Lys Val Thr Asp Glu Gly Glu Gly Gly Asn Gln Ile

245

Val Gln Tyr Leu Val Ala Gln Lys Ile Ile

250

<211> 30	
<212> DNA	
<213> Mycobacterium tuberculosis	
•	
<400> 159	
gcttccatgg gcgaggcaca ggcgtgggaa	30
gereeargy gegaggeaca ggegegggaa	30
010 160	
<210> 160	
<211> 30	
<212> DNA	
<213> Mycobacterium tuberculosis	
<400> 160	
ctgagatcta gaatgccaca gggaactgtg	30
<210> 161	
<211> 30	
<212> DNA	
<213> Mycobacterium tuberculosis	
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<400> 161	
	30
tetecegggg gtaacteaga gegageggae	30
<210> 162	
<211> 27	
<212> DNA	
<213> Mycobacterium tuberculosis	
<400> 162	
ctgagatcta tgaacgtcac cgtatcc	27
<210> 163	
<211> 27	
<212> DNA	
<213> Mycobacterium tuberculosis	
<400> 163	
tetecegggg eteacecace ggecacg	27
5555	
<210> 164	
<211> 30	
<211> 30 <212> DNA	
<213> Mycobacterium tuberculosis	
(213) MyCobaccerium cuberculosis	
400. 164	
<400> 164	
ctgagatcta tggcaacacg ttttatgacg	30
<210> 165	

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<210> 170
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<212> PRT
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<220>
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                  5
                                     10
<210> 171
<211> 15
<212> PRT
<213> Mycobacterium tuberculosis
<400> 171
Pro Lys Arg Ser Glu Tyr Arg Gln Gly Thr Pro Asn Trp Val Asp
<210> 172
<211> 404
<212> PRT
<213> Mycobacterium tuberculosis
<400> 172
Met Ala Thr Val Asn Arg Ser Arg His His His His His His His
 1
                  5
                                     10
                                                          15
Ile Glu Gly Arg Ser Phe Ser Arg Pro Gly Leu Pro Val Glu Tyr Leu
                                                     30
             20
                                 25
Gln Val Pro Ser Pro Ser Met Gly Arg Asp Ile Lys Val Gln Phe Gln
                             40
         35
Ser Gly Gly Asn Asn Ser Pro Ala Val Tyr Leu Leu Asp Gly Leu Arg
     50
                         55
Ala Gln Asp Asp Tyr Asn Gly Trp Asp Ile Asn Thr Pro Ala Phe Glu
                     70
                                         75
Trp Tyr Tyr Gln Ser Gly Leu Ser Ile Val Met Pro Val Gly Gln
```

Ser	Ser	Phe	Tyr	Ser	Asp	Trp	Tyr	Ser	Pro	Ala	Cys	Gly	Lys	Ala	Gly
			100					105					110		

- Cys Gln Thr Tyr Lys Trp Glu Thr Phe Leu Thr Ser Glu Leu Pro Gln
 115 120 125
- Trp Leu Ser Ala Asn Arg Ala Val Lys Pro Thr Gly Ser Ala Ala Ile 130 135 140
- Pro Gln Gln Phe Ile Tyr Ala Gly Ser Leu Ser Ala Leu Leu Asp Pro 165 170 175
- Ser Gln Gly Met Gly Pro Ser Leu Ile Gly Leu Ala Met Gly Asp Ala 180 185 190
- Gly Gly Tyr Lys Ala Ala Asp Met Trp Gly Pro Ser Ser Asp Pro Ala 195 200 205
- Trp Glu Arg Asn Asp Pro Thr Gln Gln Ile Pro Lys Leu Val Ala Asn 210 215 220
- Asn Thr Arg Leu Trp Val Tyr Cys Gly Asn Gly Thr Pro Asn Glu Leu 225 230 235 240
- Gly Gly Ala Asn Ile Pro Ala Glu Phe Leu Glu Asn Phe Val Arg Ser 245 250 255
- Ser Asn Leu Lys Phe Gln Asp Ala Tyr Asn Ala Ala Gly Gly His Asn 260 265 270
- Ala Val Phe Asn Phe Pro Pro Asn Gly Thr His Ser Trp Glu Tyr Trp
 275 280 285
- Gly Ala Gln Leu Asn Ala Met Lys Gly Asp Leu Gln Ser Ser Leu Gly 290 295 300
- Ala Gly Lys Leu Ala Met Thr Glu Gln Gln Trp Asn Phe Ala Gly Ile 305 310 315 320
- Glu Ala Ala Ser Ala Ile Gln Gly Asn Val Thr Ser Ile His Ser 325 330 335
- Leu Leu Asp Glu Gly Lys Gln Ser Leu Thr Lys Leu Ala Ala Ala Trp 340 345 350

Arg Ala Gln Asp Asp Tyr Asn Gly Trp Asp Ile Asn Thr Pro Ala Phe 165 170 175

Glu Trp Tyr Tyr Gln Ser Gly Leu Ser Ile Val Met Pro Val Gly Gly
180 185 190

Gln Ser Ser Phe Tyr Ser Asp Trp Tyr Ser Pro Ala Cys Gly Lys Ala 195 200 205

Gly Cys Gln Thr Tyr Lys Trp Glu Thr Phe Leu Thr Ser Glu Leu Pro 210 215 220

Gln Trp Leu Ser Ala Asn Arg Ala Val Lys Pro Thr Gly Ser Ala Ala 225 230 235 240

Ile Gly Leu Ser Met Ala Gly Ser Ser Ala Met Ile Leu Ala Ala Tyr
245 250 255

His Pro Gln Gln Phe Ile Tyr Ala Gly Ser Leu Ser Ala Leu Leu Asp 260 265 270

Pro Ser Gln Gly Met Gly Pro Ser Leu Ile Gly Leu Ala Met Gly Asp 275 280 285

Ala Gly Gly Tyr Lys Ala Ala Asp Met Trp Gly Pro Ser Ser Asp Pro 290 295 300

Ala Trp Glu Arg Asn Asp Pro Thr Gln Gln Ile Pro Lys Leu Val Ala 305 310 315 320

Asn Asn Thr Arg Leu Trp Val Tyr Cys Gly Asn Gly Thr Pro Asn Glu 325 330 335

Leu Gly Gly Ala Asn Ile Pro Ala Glu Phe Leu Glu Asn Phe Val Arg
340 345 350

Ser Ser Asn Leu Lys Phe Gln Asp Ala Tyr Asn Ala Ala Gly Gly His 355 360 365

Asn Ala Val Phe Asn Phe Pro Pro Asn Gly Thr His Ser Trp Glu Tyr 370 375 380

Trp Gly Ala Gln Leu Asn Ala Met Lys Gly Asp Leu Gln Ser Ser Leu 385 390 395 400

Gly Ala Gly

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<210> 174
<211> 291
<212> DNA
<213> Mycobacterium tuberculosis
<400> 174
atgtcgcaga ttatgtacaa ctatccggcg atgatggctc atgccgggga catggccggt 60
tatgegggea egetgeagag ettgggggee gatategeea gtgageagge egtgetgtee 120
agtgcttggc agggtgatac cgggatcacg tatcagggct ggcagaccca gtggaaccag 180
gccctagagg atctggtgcg ggcctatcag tcgatgtctg gcacccatga gtccaacacc 240
atggcgatgt tggctcgaga tggggccgaa gccgccaagt ggggcggcta g
                                                                   291
<210> 175
<211> 96
<212> PRT
<213> Mycobacterium tuberculosis
<400> 175
Met Ser Gln Ile Met Tyr Asn Tyr Pro Ala Met Met Ala His Ala Gly
                                     10
                  5
Asp Met Ala Gly Tyr Ala Gly Thr Leu Gln Ser Leu Gly Ala Asp Ile
             20
                                 25
Ala Ser Glu Gln Ala Val Leu Ser Ser Ala Trp Gln Gly Asp Thr Gly
         35
                             40
Ile Thr Tyr Gln Gly Trp Gln Thr Gln Trp Asn Gln Ala Leu Glu Asp
Leu Val Arg Ala Tyr Gln Ser Met Ser Gly Thr His Glu Ser Asn Thr
                     70
                                         75
Met Ala Met Leu Ala Arg Asp Gly Ala Glu Ala Ala Lys Trp Gly Gly
```

<210> 176

<211> 363

<212> DNA

<213> Mycobacterium tuberculosis

85

90

<400> 176
gtgtcgcaga gtatgtacag ctacceggcg atgacggca atgtcggaga catggccggt 60
tatacgggca cgacgcagag cttgggggcc gatatcgcca gtgagcgcac cgcgccgtcg 120
cgtgcttgcc aaggtgatct cgggatgagt catcaggact ggcaggccca gtggaatcag 180
gccatggagg ctctcgcgg ggcctaccgt cggtgccggc gagcactacg ccagatcggg 240
gtgctggaaa ggccggtagg cgattcgtca gactgcggaa cgattagggt ggggtcgttc 300
cggggtcggt ggctggaccc gcgccatgcg ggtccagcca cggccgcga cgccggagac 360
taa

<210> 177

<211> 120

<212> PRT

<213> Mycobacterium tuberculosis

<400> 177

Met Ser Gln Ser Met Tyr Ser Tyr Pro Ala Met Thr Ala Asn Val Gly $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asp Met Ala Gly Tyr Thr Gly Thr Thr Gln Ser Leu Gly Ala Asp Ile 20 25 30

Ala Ser Glu Arg Thr Ala Pro Ser Arg Ala Cys Gln Gly Asp Leu Gly
35 40 45

Met Ser His Gln Asp Trp Gln Ala Gln Trp Asn Gln Ala Met Glu Ala 50 55 60

Leu Ala Arg Ala Tyr Arg Arg Cys Arg Arg Ala Leu Arg Gln Ile Gly 65 70 75 80

Val Leu Glu Arg Pro Val Gly Asp Ser Ser Asp Cys Gly Thr Ile Arg 85 90 95

Val Gly Ser Phe Arg Gly Arg Trp Leu Asp Pro Arg His Ala Gly Pro
100 105 110

Ala Thr Ala Ala Asp Ala Gly Asp 115 120

<210> 178

<211> 297

<212> DNA

<213> Mycobacterium tuberculosis

<400> 178

atggcctcgc gttttatgac ggatccgcac gcgatgcggg acatggcggg ccgttttgag 60 gtgcacgccc agacggtgga ggacgaggct cgccggatgt gggcgtccgc gcaaaacatc 120

tegggegeg getggagtgg catggeegag gegacetege tagacaccat gacecagatg 180 aateaggegt ttegcaacat egtgaacatg etgeaeggg tgegtgaegg getggttege 240 gaegeeaaca actaegaaca geaagageag geeteeeage agateeteag eagetga 297

<210> 179

<211> 98

<212> PRT

<213> Mycobacterium tuberculosis

<400> 179

Met Ala Ser Arg Phe Met Thr Asp Pro His Ala Met Arg Asp Met Ala 1 5 10 15

Gly Arg Phe Glu Val His Ala Gln Thr Val Glu Asp Glu Ala Arg Arg
20 25 30

Met Trp Ala Ser Ala Gln Asn Ile Ser Gly Ala Gly Trp Ser Gly Met
35 40 45

Ala Glu Ala Thr Ser Leu Asp Thr Met Thr Gln Met Asn Gln Ala Phe 50 55 60

Arg Asn Ile Val Asn Met Leu His Gly Val Arg Asp Gly Leu Val Arg 65 70 75 80

Asp Ala Asn Asn Tyr Glu Gln Glu Gln Ala Ser Gln Gln Ile Leu 85 90 95

Ser Ser

<210> 180

<211> 297

<212> DNA

<213> Mycobacterium tuberculosis

<400> 180

atggeeteac gttttatgac ggateegeac gegatgeggg acatggeggg cegttttgag 60 gtgeacgeec agacggtgga ggaegagget egeeggatgt gggegteege geaaaacatt 120 teeggtgegg getggagtgg catggeegag gegacetege tagacaceat ggeecagatg 180 aateaggegt ttegeaacat egtgaacatg etgeacggg tgegtgaegg getggttege 240 gaegeeaaca actaegagea geaagageag geeteecage agateeteag eagetaa 297

<210> 181

<211> 98

<212> PRT

<213> Mycobacterium tuberculosis

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<400> 181
Met Ala Ser Arg Phe Met Thr Asp Pro His Ala Met Arg Asp Met Ala
                                      10
Gly Arg Phe Glu Val His Ala Gln Thr Val Glu Asp Glu Ala Arg Arg
             20
                                 25
Met Trp Ala Ser Ala Gln Asn Ile Ser Gly Ala Gly Trp Ser Gly Met
                             40
Ala Glu Ala Thr Ser Leu Asp Thr Met Ala Gln Met Asn Gln Ala Phe
                         55
Arg Asn Ile Val Asn Met Leu His Gly Val Arg Asp Gly Leu Val Arg
                     70
                                         75
Asp Ala Asn Asn Tyr Glu Gln Glu Gln Ala Ser Gln Gln Ile Leu
                 85
                                     90
Ser Ser
<210> 182
<211> 297
<212> DNA
<213> Mycobacterium tuberculosis
<400> 182
atggcctcac gttttatgac ggatccgcat gcgatgcggg acatggcggg ccgttttgag 60
gtgcacgccc agacggtgga ggacgaggct cgccggatgt gggcgtccgc gcaaaacatt 120
teeggtgegg getggagtgg catggeegag gegacetege tagacaccat gacetagatg 180
aatcaggegt ttegeaacat egtgaacatg etgeaegggg tgegtgaegg getggttege 240
gacgccaaca actacgaaca gcaagagcag gcctcccagc agatcctgag cagctag
<210> 183
<211> 98
<212> PRT
<213> Mycobacterium tuberculosis
<400> 183
Met Ala Ser Arg Phe Met Thr Asp Pro His Ala Met Arg Asp Met Ala
                  5
                                     10
Gly Arg Phe Glu Val His Ala Gln Thr Val Glu Asp Glu Ala Arg Arg
             20
                                 25
                                                     30
```

Met Trp Ala Ser Ala Gln Asn Ile Ser Gly Ala Gly Trp Ser Gly Met 45 40 35 Ala Glu Ala Thr Ser Leu Asp Thr Met Thr Gln Met Asn Gln Ala Phe 50 55 60 Arg Asn Ile Val Asn Met Leu His Gly Val Arg Asp Gly Leu Val Arg 80 65 70 75 Asp Ala Asn Asn Tyr Glu Gln Glu Gln Ala Ser Gln Gln Ile Leu 85 90 95 Ser Ser <210> 184 <211> 297 <212> DNA <213> Mycobacterium tuberculosis <400> 184 atgacetege gttttatgae ggateegeae gegatgeggg acatggeggg cegttttgag 60 gtgcacgccc agacggtgga ggacgaggct cgccggatgt gggcgtccgc gcaaaacatt 120 teeggegegg getggagtgg catggeegag gegaeetege tagacaccat gaeecagatg 180 aatcaggegt ttegeaacat egtgaacatg etgeaegggg tgegtgaegg getggttege 240 qacqccaaca actacqaaca gcaagagcag gcctcccagc agatcctcag cagctga <210> 185 <211> 98 <212> PRT <213> Mycobacterium tuberculosis <400> 185 Met Thr Ser Arg Phe Met Thr Asp Pro His Ala Met Arg Asp Met Ala 15 1 5 10 Gly Arg Phe Glu Val His Ala Gln Thr Val Glu Asp Glu Ala Arg Arg 20 30 Met Trp Ala Ser Ala Gln Asn Ile Ser Gly Ala Gly Trp Ser Gly Met 45 35 40 Ala Glu Ala Thr Ser Leu Asp Thr Met Thr Gln Met Asn Gln Ala Phe 50 55 60 Arg Asn Ile Val Asn Met Leu His Gly Val Arg Asp Gly Leu Val Arg 80 70 75 65

Asp Ala Asn Asn Tyr Glu Gln Gln Glu Gln Ala Ser Gln Gln Ile Leu 85 90 95

Ser Ser

<210> 186	
<211> 20	
<212> DNA	
<213> Mycobacterium tuberculosis	
<400> 186	
ggaatgaaaa ggggtttgtg	20
<210> 187	
<211> 20	
<212> DNA	
<213> Mycobacterium tuberculosis	
<400> 187	
gaccacgccc gcgccgtgtg	20
<210> 188	
<211> 27	
<212> DNA	
<213> Mycobacterium tuberculosis	
<400> 188	
gcaacacccg ggatgtcgca gattatg	27
<210> 189	
<211> 30	
<212> DNA	
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5	мес 1	ser	GTII	тте	ме с 5	тАт	ASII	тАт	PIO	10	Met	пеи	GTĀ	птъ	15	GTA	
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-10	Mec	50	1113	GIII	лър	112	55	ALG	0111	110	ADI	60	mu	1100	GIU	1114	
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45	ьец 65	Ala	Arg	АТА	TÄT	70	Arg	Cys	Arg	Arg	75	Leu	Arg	GTII	тте	80	
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50	VUL	Dea	GIG	211.9	85	val	OLY	11010	DCI	90	11010	CJD	013	1111	95	211 9	
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	vaı	GΤĀ	ser	100	arg	ЭΤΆ	arg	Trp	ьеи 105	ASP	Pro	Arg	HIS	110	СΤĀ	Pro	
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	45	gcc Ala	agt Ser	gag Glu 35	Gln	gcc Ala	gtg Val	ctg Leu	tcc Ser 40	Ser	gct Ala	tgg Trp	cag Gln	ggt Gly 45	gat Asp	acc Thr	Gly	144
	50	atc Ile	acg Thr 50	Туг	cag Gln	ggc Gly	tgg Trp	cag Gln 55	Thr	cag Gln	tgg Trp	aac Asn	cag Gln 60	Ala	cta Leu	ı gag ı Glu	gat Asp	192
	55	ctg Leu	Va1	g cgg Arg	gcc Ala	tat Tyr	cag Glr	ı Ser	atç Met	tct Ser	ggc Gly	acc Thr	His	gag Glu	tco Ser	aac Asr	acc Thr	240

Hard Hard

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atg geg atg ttg get ega gat ggg gee gaa gee gee aag tgg gge gge Met Ala Met Leu Ala Arg Asp Gly Ala Glu Ala Ala Lys Trp Gly Gly 90

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    Ile Thr Tyr Gln Gly Trp Gln Thr Gln Trp Asn Gln Ala Leu Glu Asp
20
                             55
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                                                  10
                                                                                      60
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10		<2 <2		60 DNA	obact	teri	ım tı	ıbero	culos	sis								
1 5		<2 <2		(1)	(6	50)												
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25	cgg Arg	_		-														60
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The Cont. Unit

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                                                                     15
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         Asp Gly Ala Glu
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ļ.a
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That I I I I'm Kit hat hat took
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         Asp Met
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          Ser Leu
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          Tyr Ala Gly Thr Leu Gln Ser Leu Gly Ala Glu Ile Ala Val Glu Gln
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                            5
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          cgg
          Arg
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South Carle Bash of High Hards April Bash
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          Arg
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     50 Ser Leu Gly Ala
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                                                                                                                                                        10
                               1
IJ
                              Thr Ala Pro Ser
H...H
                                                                       20
1
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                45
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                50 Asp Ile Ala Ser Glu Arg Thr Ala Pro Ser Arg Ala Cys Gln Gly Asp
                                  1
                                                                                                                                                           10
                              ctc ggg atg agt
                                                                                                                                                                                                                                                                            60
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         cgt gct tgc caa ggt gat ctc ggg atg agt cat cag gac tgg cag gcc
                                                                                  48
         Arg Ala Cys Gln Gly Asp Leu Gly Met Ser His Gln Asp Trp Gln Ala
     25
                                                10
         1
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         cag tgg aat cag
# T. ...
         Gln Trp Asn Gln
30
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35
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         Arg Ala Cys Gln Gly Asp Leu Gly Met Ser His Gln Asp Trp Gln Ala
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    15
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                                                                                   48
         Ala Met Glu Ala Leu Ala Arg Ala Tyr Arg Arg Cys Arg Arg Ala Leu
     30
                                                                                   60
         cgc cag atc ggg
£į.
         Arg Gln Ile Gly
firm 2007
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dini.
                <210> 249
13
                <211> 20
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     40
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          Ala Met Glu Ala Leu Ala Arg Ala Tyr Arg Arg Cys Arg Arg Ala Leu
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                                         10
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                 20
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           <210> 254
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           <212> DNA
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55
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                     5
                                          10
                                                                            60
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    Pro Arg His Ala
                20
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                                         10
                                                             15
    Asp Ala Gly Asp
                20
```

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